

SCORE Search Results Details for Application 10777592 and Search Result us-10-777-592- 1.rng.

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This page gives you Search Results detail for the Application 10777592 and Search Result us-10-777-592-1.rng.

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GenCore version 5.1.9
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1-1

rng

RT prim

OM nucleic - nucleic search, using sw model

Run on: August 17, 2006, 21:35:34 ; Search time 165.5 Seconds
(without alignments)
505.541 Million cell updates/sec

Title: US-10-777-592-1
Perfect score: 12
Sequence: 1 gttacagcacag 12

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 5244920 seqs, 3486124231 residues

Total number of hits satisfying chosen parameters: 10489840

Minimum DB seq length: 0
Maximum DB seq length: 20000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1000 summaries

abase : N_Geneseq_8:*

- 1: geneseqn1980s:*
- 2: geneseqn1990s:*
- 3: geneseqn2000s:*
- 4: geneseqn2001as:*
- 5: geneseqn2001bs:*
- 6: geneseqn2002as:*
- 7: geneseqn2002bs:*
- 8: geneseqn2003as:*
- 9: geneseqn2003bs:*
- 10: geneseqn2003cs:*
- 11: geneseqn2003ds:*
- 12: geneseqn2004as:*
- 13: geneseqn2004bs:*
- 14: geneseqn2005s:*
- 15: geneseqn2006s:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result	No.	Score	Query Match	Length	DB	ID	Description
	1	12	100.0	18	3	AAA55650	Aaa55650 TRAF5 ant
c	2	12	100.0	24	2	AAV01735	Aav01735 PCR prime
	3	12	100.0	27	12	ADJ14698	Adj14698 Debrisoqu
	4	12	100.0	27	12	ADJ14710	Adj14710 Debrisoqu
	5	12	100.0	27	12	ADO60813	Ado60813 Human deb
	6	12	100.0	27	12	ADO60800	Ado60800 Human deb
	7	12	100.0	27	12	ADO61063	Ado61063 Human deb
	8	12	100.0	27	14	AEC90233	Aec90233 CYP2D6 ge
	9	12	100.0	27	14	AEC89964	Aec89964 CYP2D6 ge
	10	12	100.0	27	14	AEC90227	Aec90227 CYP2D6 ge
	11	12	100.0	27	14	AEC89977	Aec89977 CYP2D6 ge
c	12	12	100.0	97	4	AAI26366	Aai26366 Probe #16
c	13	12	100.0	97	4	ABA74237	Aba74237 Human foe
c	14	12	100.0	97	4	AAI54694	Aai54694 Probe #23
c	15	12	100.0	97	4	ABA39221	Aba39221 Probe #17
c	16	12	100.0	97	4	AAK48865	Aak48865 Human bon
c	17	12	100.0	97	4	AAK22698	Aak22698 Human bra
c	18	12	100.0	97	4	ABS48536	Abs48536 Human liv
c	19	12	100.0	97	6	ABS22555	Abs22555 Human gen
	20	12	100.0	162	14	ADW83986	Adw83986 MAP3K9 ma
c	21	12	100.0	165	2	AAT22527	Aat22527 Human gen
c	22	12	100.0	190	10	ACD98264	Acd98264 Human col
c	23	12	100.0	198	14	AEB50463	Aeb50463 Human myo
c	24	12	100.0	254	10	ACD95327	Acd95327 Human col
	25	12	100.0	279	8	ABX36714	Abx36714 Bovine ES
c	26	12	100.0	279	8	ABX38879	Abx38879 Bovine ES
c	27	12	100.0	288	2	AAT26835	Aat26835 Human gen
	28	12	100.0	297	8	ABX46205	Abx46205 Bovine ES
	29	12	100.0	307	10	ADF80624	Adf80624 Leukaemia
c	30	12	100.0	321	2	AAV88302	Aav88302 EST clone
c	31	12	100.0	325	8	ABX36123	Abx36123 Bovine ES
	32	12	100.0	331	14	ADW83983	Adw83983 MAP3K9 ma
c	33	12	100.0	351	3	AAC14703	Aac14703 Human sec
c	34	12	100.0	367	8	ABX44174	Abx44174 Bovine ES
c	35	12	100.0	374	4	AAS50222	Aas50222 Staphyloc
	36	12	100.0	374	8	ACA17436	Aca17436 Prokaryot
	37	12	100.0	394	10	ADE29044	Ade29044 Human UBE
	38	12	100.0	396	14	AEC23319	Aec23319 Human myo
	39	12	100.0	396	15	AEF72645	Aef72645 Myocardia
	40	12	100.0	402	5	AAF64558	Aaf64558 Novel hum
c	41	12	100.0	403	6	ABL82486	Abl82486 Human ova
	42	12	100.0	408	5	AAF66062	Aaf66062 Novel hum
c	43	12	100.0	410	8	ABX42193	Abx42193 Bovine ES
c	44	12	100.0	426	8	ABX37493	Abx37493 Bovine ES
	45	12	100.0	428	5	AAF64893	Aaf64893 Novel hum
	46	12	100.0	428	14	ADY66531	Ady66531 S. manson
	47	12	100.0	442	6	ABN61747	Abn61747 Human can
	48	12	100.0	443	6	ABT07032	Abt07032 Human ova
	49	12	100.0	443	8	ABX72910	Abx72910 Human ova
c	50	12	100.0	445	13	ADQ58217	Adq58217 Novel can
	51	12	100.0	457	2	AAV23759	Aav23759 Pac gene
c	52	12	100.0	459	6	ABL84710	Abl84710 Human ova